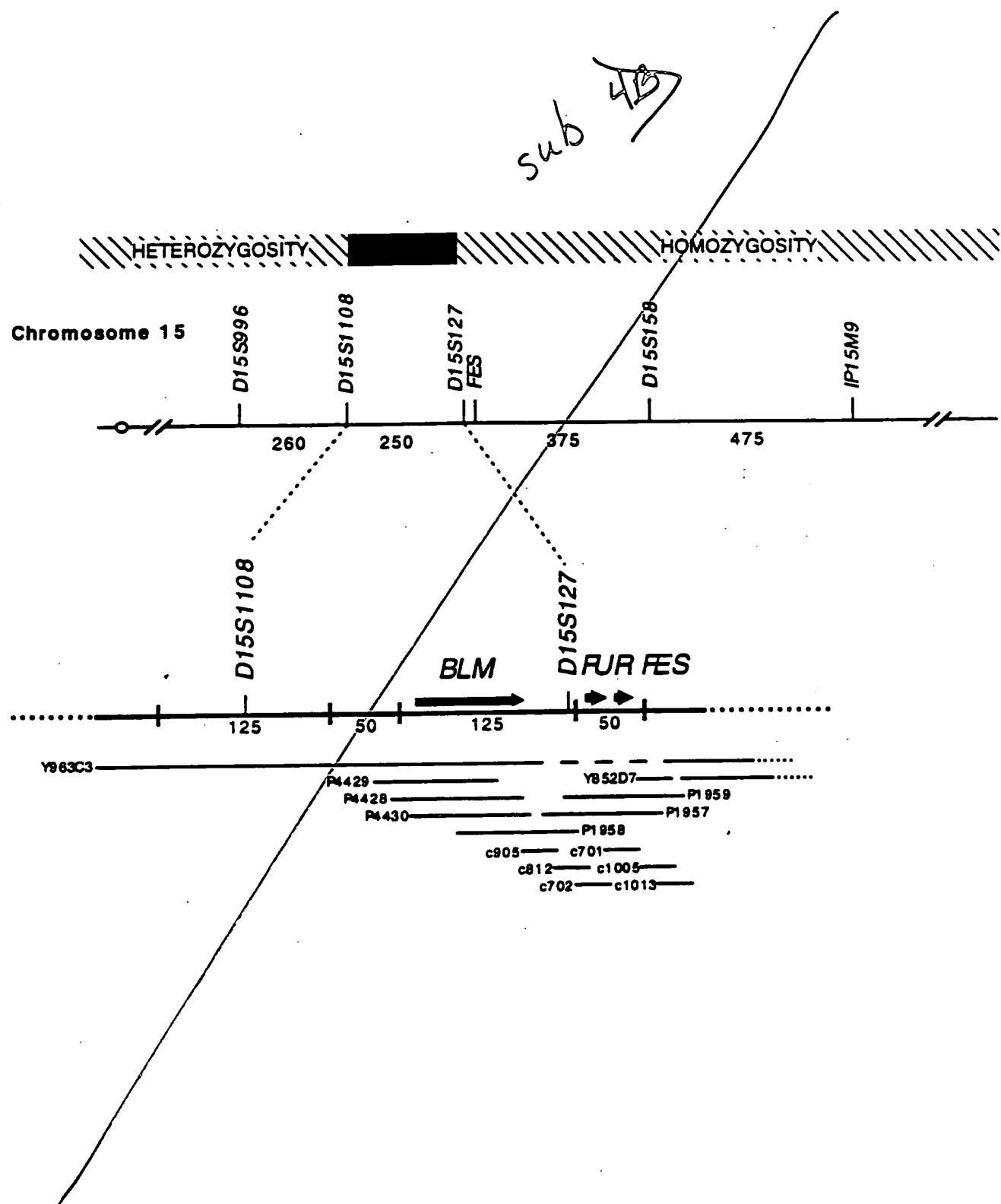


FIGURE 1A

○ 〔 〕 〔 〕 〔 〕 〔 〕 〔 〕 〔 〕 〔 〕 〔 〕 〔 〕 〔 〕

FIGURE 1B

D15S1108 PF PH PJ PM
11(LaTH)


<u>59(FrFit)</u>	PF	P ^h	P ^d	RM
<u>87(AIFra)</u>	PF	P ^h	P ^d	RM

NR8(KeS01)

D15S127 PF PH P1 P1 PM

PF P^h P^f PW PF P^h P^f PW

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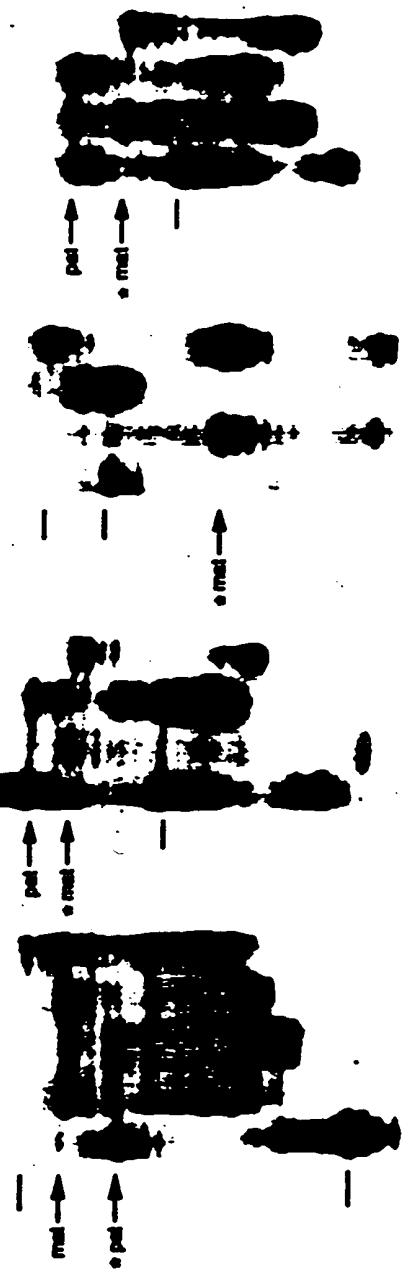


FIGURE 2

FIGURE 2 (cont'd)

TGCGGACATGATTTCTCAAGATTACAAAAGAATGAATATGCCAGAAGTTCCCTCTGTTCCGGTGAATGGCTCT 2560
803 W G H D F R Q D Y K R M N M L R Q K F P S V P V M A L
TACGGCCACAGCTAAATCCCAGGGTACAGAAGGACTCTGACTCAGCTGAAGATTCAGACCTCAGGTGTTACCATGA 2640
830 T A T A N P R V Q K D I L T Q L K I L R P Q V F S M
GCTTTAACAGACATAATCTGAAATACTATGTATTACCGAAAAGCCTAAAAGGTGGCATTGATTCAGCTAGAATGGATC 2720
856 S F N R H N L K Y Y V L P K K V A F D C L E W I
AGAAAGCACCACCCATATGATTCAAGGATAATTACTGCTCTCCAGGGAGAAATGTGACACCATGGCTGACACGGTACA 2800
883 R K H H P Y D S G I I Y C L S R R E C D T M A D T L Q
GAGAGATGGGCTCGCTGCTTGCCTACCATGCTGGCTCAGTGAATCTGCCAGAGATGAAGTGCAGCAGAAGTGGATTA 2880
910 R D G L A A L A Y H A G L S D S A R D E V Q Q K W I
ATCAGGATGGCTGCTGAGGTATCTGCTACAATTGCAATTGGAAATGGGATTGACAAACCGACGTGGGATTTGATTC 2960
936 N Q D G C Q V I C A T I A F G M G I D K P D V R F V I
CATGGATCTCTCCCTAAATCTGAGGGGTTACTACCAAGAATCTGGCAGAGCTGGAAAGATGGAAATATCTCACTG 3040
963 H A S L P K S V E G Y Y Q E S G R A G R D G E I S H C
CCTGCTTTCTATACCTATCATGATGTGACAGACTGAAAAGACTTATAATGATGGAAAAGATGGAAACCATCATAAA 3120
990 L L F Y T Y H D V T R L K R L I M M E K D G N H H T
GAGAAACTCACTTCATAATTGATAGCATGGTACATTACTGTGAAAATATAACCGAATGCCAGGAGAATACAGCTTTG 3200
1016 R E T H F N N L Y S M V H Y C E N I T E C R R I Q L L
GCCTACTTTGGTGAAGATTTAAATCTGATTTTCTGAAACACCCAGATGTTCTGTGATAATTGCTGAAAAAC 3280
1043 A Y F G E N G F N P D F C K K H P D V S C D N C C K T
AAAGGATTATAAAACAAGAGATGTGACTGACGATGTGAAAAGTATTGTAAGATTGTTCAAGAACATAGTCATCACAAG 3360
1070 K D Y K T R D V T D D V K S I V R F V Q E H S S S Q
GAATGAGAAAATATAAACATGTAGGTCTCTGGAAAGATTTACTATGAATATGCTGGCGACATTCTGGGAGTAAG 3440
1106 G M R N I K H V G P S G R F T M N M L V D I F L G S K
ACTGCAAAATCCAGTCAGTATATTGAAAAGGATCTGCTTATTCAGCACACAATGCCAGAGACTTTAAAAGCT 3520
1123 S A K I Q S G I F G K G S A Y S R H N A E R L F K K L
GAATCTGACAAGATTTGGATGAAGACTTATATATCAATGCCAATGACCAGGGGATGCTTATGTGATGCTGGAAAATA 3600
1150 J L D K I L D E D L Y I N A N D Q A I A Y V M L G N
AAACCAAAACTGTACTAAATGCCAATTAAAGGTAGACTTTATGGAAACAGAAAATCCACAGTGTGAAAAACAAAAA 3680
1176 K I Q T V L N G N L K V D F M E T E N S S S V K K Q K
GCCTTAGTAGCAAAGTGTCTCAGAGGGAAAGAGATGGTAAAGATGGAGAACTTACAGAAGTCTGCAAATCTCT 3760
1203 A L V A K V S Q R E E M V K K C L G E L T E V C K S L
GGGGAAAGTTTTGGTGTCCATTACTTCATAATTAAACCGTCACTCTCAAGAACCTTGCAAGAATCTTATCTCTG 3840
1230 G K V F G V H Y F N I F N T V T L K K L A E S L S S
ATCCTGAGGTTTGCTTCAATTGATGGTGTACTGAAAGACAACTGGAAAAATATGGTGGGAAAGTGAATTCACTGATTA 3920
1256 D P E V L L Q I D G V T E D K L E K Y G A E V I S V L
CAGAAATACTCTGAATGGACATGCCAGCTGAAGACAGTCCCCAGGGATAAGCTGCTCCAGCACAGAGGGCCCCGGAG 4000
1283 Q K Y S E W T S P A E D S S P G I S L L S S S R G P G R
AACTGCGGCTGAGGAGCTTGACGGAGAAATACCGTATCTCCACTACTTGTCAAGTAAAACCAGAAAATGAAAGGAAGA 4080
1310 S A A E E L D E E I P V S S H Y F A S K T R N E R K
GGAAAAAGATGCCAGCTCCAAAGGTCTAAGAGGAGAAAATCTGCTCCAGTGGTCCAGGCAAGGCAAAGGGGGGCTGCC 4160
1336 R K K M P A S Q R S K R R K T A S S G S K A K G G S A
ACATGTAGAAAAGATATCTTCCAAAACGAAATCTCCACCATGGATCCAGGTCAAGCTCACATACTTCTCAAGCGAC 4240
1363 T C R K I S S K T K S S S I I G S S S A S H T S Q A T
ATCAGGAGCCAATAGCAAATTGGGATTATGGCTCCACCGAAGCTATAAAATAGACCGTTCTTAAGCTTCATATGCAT 4320
1390 S G A N S K L G I M A P P K P I N R P F L K P S Y A
TCTCATAAcaaccgaatctcaatgtacatagaccctttttttgtcagcatctgaccatctgtgactataaagctg 4400
1416 F S .
ttatttcttgttatccaaaaaaaaaaaaaaaaaaaaaa 4437

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FIGURE 3

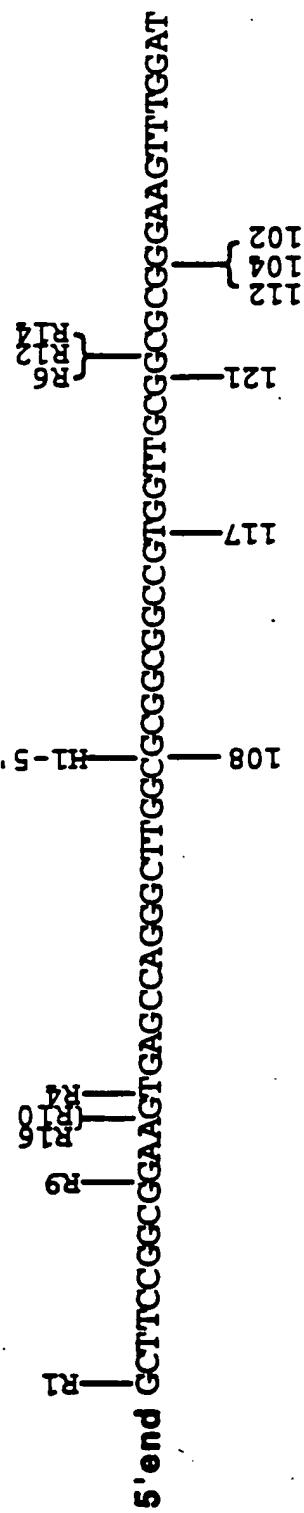


FIGURE 4

I

Ia

649 F E H T K E M M K I F K K C G L E M I N E R T N Q V A M A L G E D N F I E M P T G G G K S I C Y O L P A C V - - - - - T V W H S P L R S I T V D Q V BLM
 74 F P W S G K V K D I L Q N V E K C E K P E R I Q L E T I N V I M A G E V P L V M P T G G G K S I C Y O L P A C C - - - - - S A I F T L V I C P L I S L M E D O L R E Q L
 659 Y E W S D E V L Y R L E E V E K U P G E R P A N O L E A V N A T L O G U V F V I M P T G G G K S I C Y O L P A V V K S G K T H G T I V I S P I I S L M O D O V SGS1
 16 -----V D O E T R G Y Q O O E R P G H E E I E D T V S Q R D L V M P T G G G K S I C Y O L P A L L - - - - - L N G L E W V S P I I S L M Q D O V recQ

II

725 Q K L T S L D E P A T Y E T G D K Y D S E A T N I Y L Q L S K E D P I E K L L Y V I P E K I C A S N R G I S T E E N C Y E R K L I A R F I D E A H C V S Q W G BLM
 150 M V I K Q L G I S A T M I N A S S K E H V K W I H D E M V N K N S E L K I T V V T P E K I A K S K M F M S R L E K A Y E A R R F T R I A V D E V H C C S Q W G R E Q L
 739 E H L L N K N J I K S M F S S R G I A E Q R Q T F N L F I N - - E L N D E V Y S P E M T S A S E O C K R A I S R L Y P D G K C A R I D V D E A H C V S N W G SGS1
 83 D Q A Q A N G V A S A C N S T O P E Q O L E J M T - - G C R T G S R U L Y L A P E R L - - - E L D N F I E H G - - H W N P V I L L A V D E A H C I S Q W G recQ

III

805 H D E R Q D Y X R M M I H O K P E S V E V I A L T A T A N P R Y Q D I L T O E K I L R P O V E S M S E N R H N I C Y V I P K P K K V A - - E D C L E W BLM
 230 H D F R E D Y K A L G I A K R O E P N A S L I G L T T A T A T N H M L T D A Q K I E C E K C F T S T A S E N H P N L - - E E V R Q K P S T E D F I E D I V K L R E Q L
 817 H D F R E D Y K E L K F F K R E Y E D I E M I A L T A T A S E Q V R M D I L H M L E K E P V P L K O S E N R T N L E V E N K T K T N T - - E E I C D A SGS1
 157 H D E R P E Y A A L G Q L E O R E P T L F E A L T A T D I T T S O D I E V R L G I N D B L I Q I S S E D R P N I R Y - M I M E K F K P L D Q L M - - - R Y recQ

IV

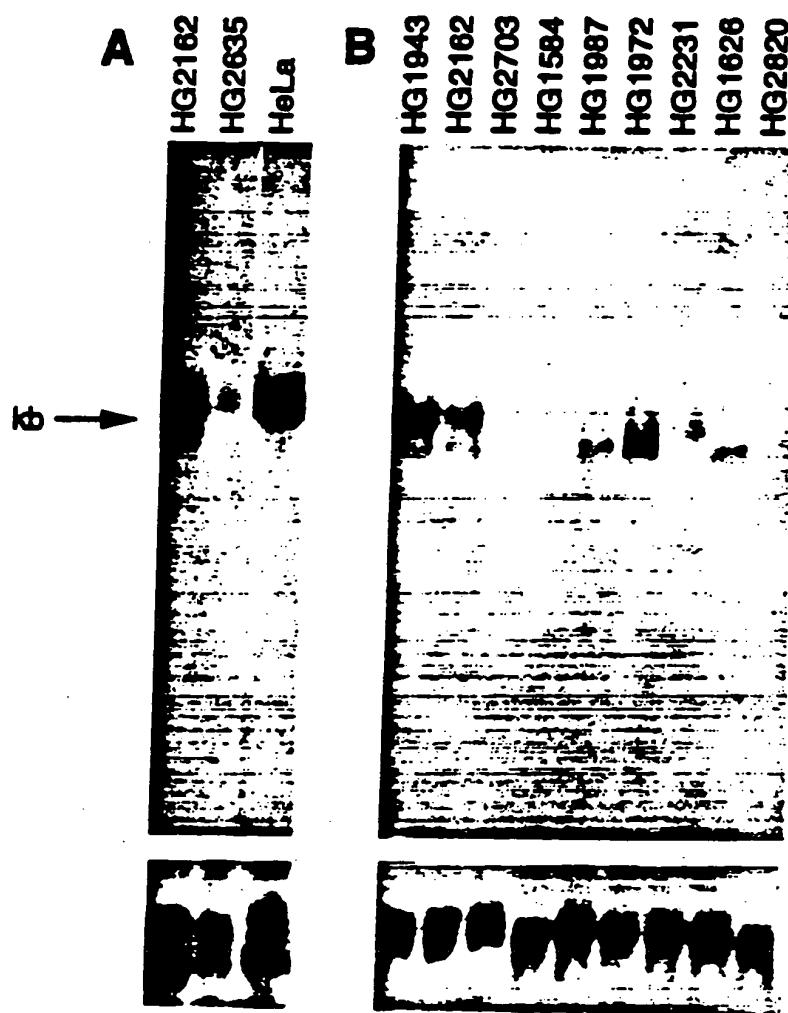
882 E R K H H P Y D S G T I Y C L S R R E G D T M A D T L O E D G L A A L A Y H A G I S D S A R D E V O Q W I N O D C C O V I D A T I A F G M G I D K P D V R F V BLM
 309 E N G R Y K G Q S G I L Y C F S Q I D S E Q V T V S L Q N L G I E H A G A Y H A N L E P E D K T T V H R I W S A N E - - L O I V V A T V A F G M G I D K P D V R F V R E Q L
 893 E K S R F K N Q T G I L Y C H S K K S C E O T S A Q M Q R N G I K C A Y Y H A G M E P D E R L S V O K A M O A D E - - L O I V V A T V A F G M G I D K P D V R F V SGS1
 233 Y Q E Q - R G K S G I L Y C N S R A K V E D T A A L Q S K G I S A A A Y H A G L E N N V R A D V O E F E R D D - L O I V V A T V A F G M G I N K P N V R F V recQ

V

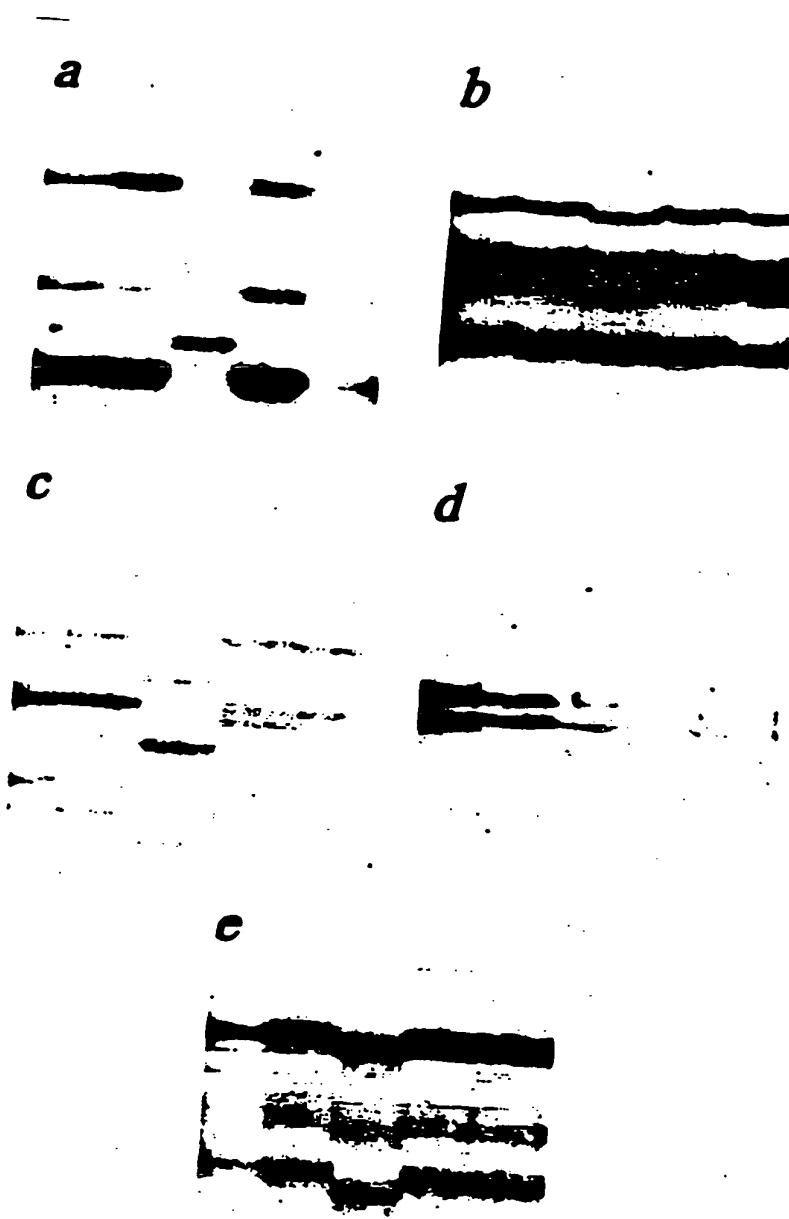
962 F I A S L P K S V E G Y Y Q E S G R A G R D G E I H C L L F Y T Y H D V T R L K R L I M M E K D G N H T R E T H F A N L Y S M V H Y C E N I T E C R R I Q E BLM
 388 F I H H M S K S M E N Y Y Q E S G R A G R D D M K A D C I L Y G E G D I F R I S S M V V M E N V G Q Q - - - - - K L Y E M V S Y C Q N I S K S R R V L M R E Q L
 972 Y H P T V F T L E G Y Q E S G R A G R D G N Y S Y C I T E F S E R D I R T M Q T M I Q K D K N L D R E N K E K H L N K L Q Q V M A Y C D N V T D C R R K G V SGS1
 314 V H F D I P H N I E S Y Y D E S G R A G R D G L P A E A M G E Y D P A D M A W I R R C L E E K P O S Q L Q D I E R H - K I N A M G A F A E A Q T - C R R L V L recQ

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FIGURES 5A and 5B



FIGURES 6A - 6E



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